```
           RAW SEQUENCE LISTING                  DATE: 05/30/2000
           PATENT APPLICATION:   US/09/079,678A   TIME: 21:32:35

           Input Set : A:\Pto.amc
           Output Set: N:\CRF3\05262000\I079678A.raw

C--> 5 (1) GENERAL INFORMATION:
     7      (i) APPLICANT: Alvarez, Vernon L.
     8                     O'Mahony, Daniel J.
     9                     Lambkin, Imelda J.
    10                     Singleton, Judith
    11                     Patterson, Catherine A.
    12                     Cagney, Gerard M.              ᵤ
    13                     Belinka, Benjamin A.
    14                     Carter, John M.
C--> 16    (ii) TITLE OF INVENTION: RANDOM PEPTIDES THAT BIND TO GASTRO-
    17                     INTESTINAL TRACT (GIT) TRANSPORT RECEPTORS AND RELATED METHODS
    20   (iii) NUMBER OF SEQUENCES: 407
    22    (iv) CORRESPONDENCE ADDRESS:
    23         (A) ADDRESSEE: Pennie & Edmonds LLP
    24         (B) STREET: 1155 Avenue of the Americas
    25         (C) CITY: New York
    26         (D) STATE: New York
    27         (E) COUNTRY: USA
    28         (F) ZIP: 10036
    30     (v) COMPUTER READABLE FORM:
    31         (A) MEDIUM TYPE: Diskette
    32         (B) COMPUTER: IBM Compatible
    33         (C) OPERATING SYSTEM: DOS
    34         (D) SOFTWARE: FastSEQ Version 2.0
C--> 36    (vi) CURRENT APPLICATION DATA:
C--> 37         (A) APPLICATION NUMBER: US/09/079,678A
C--> 38         (B) FILING DATE: 15-May-1998
    39         (C) CLASSIFICATION:
    41  (viii) ATTORNEY/AGENT INFORMATION:
    42         (A) NAME: Misrock, S. Leslie
    43         (B) REGISTRATION NUMBER: 18,872
    44         (C) REFERENCE/DOCKET NUMBER: 1101-220
    46    (ix) TELECOMMUNICATION INFORMATION:
    47         (A) TELEPHONE: 212-790-9090
    48         (B) TELEFAX: 212-869-9741
    49         (C) TELEX: 66141 PENNIE
    52 (2) INFORMATION FOR SEQ ID NO: 1:
    54      (i) SEQUENCE CHARACTERISTICS:
    55         (A) LENGTH: 44 amino acids
    56         (B) TYPE: amino acid
    57         (C) STRANDEDNESS:
    58         (D) TOPOLOGY: unknown
    60     (ii) MOLECULE TYPE: peptide
    62     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
    64  Arg Ser Gly Ala Tyr Glu Ser Pro Asp Gly Arg Gly Gly Arg Ser Tyr
    65  1               5                  10                 15
    66  Val Gly Gly Gly Gly Gly Cys Gly Asn Ile Gly Arg Lys His Asn Leu
```

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67              20                  25                  30
68  Trp Gly Leu Arg Thr Ala Ser Pro Ala Cys Trp Asp
69          35                  40
71 (2) INFORMATION FOR SEQ ID NO: 2:
73      (i) SEQUENCE CHARACTERISTICS:
74          (A) LENGTH: 44 amino acids
75          (B) TYPE: amino acid
76          (C) STRANDEDNESS:
77          (D) TOPOLOGY: unknown
79      (ii) MOLECULE TYPE: peptide
81      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
83  Ser Pro Arg Ser Phe Trp Pro Val Val Ser Arg His Glu Ser Phe Gly
84  1               5                   10                  15
85  Ile Ser Asn Tyr Leu Gly Cys Gly Tyr Arg Thr Cys Ile Ser Gly Thr
86              20                  25                  30
87  Met Thr Lys Ser Ser Pro Ile Tyr Pro Arg His Ser
88          35                  40
90 (2) INFORMATION FOR SEQ ID NO: 3:
92      (i) SEQUENCE CHARACTERISTICS:
93          (A) LENGTH: 44 amino acids
94          (B) TYPE: amino acid
95          (C) STRANDEDNESS:
96          (D) TOPOLOGY: unknown
98      (ii) MOLECULE TYPE: peptide
100     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
102 Ser Ser Ser Ser Asp Trp Gly Gly Val Pro Gly Lys Val Val Arg Glu
103 1               5                   10                  15
104 Arg Phe Lys Gly Arg Gly Cys Gly Ile Ser Ile Thr Ser Val Leu Thr
105             20                  25                  30
106 Gly Lys Pro Asn Pro Cys Pro Glu Pro Lys Ala Ala
107         35                  40
109 (2) INFORMATION FOR SEQ ID NO: 4:
111     (i) SEQUENCE CHARACTERISTICS:
112         (A) LENGTH: 44 amino acids
113         (B) TYPE: amino acid
114         (C) STRANDEDNESS:
115         (D) TOPOLOGY: unknown
117     (ii) MOLECULE TYPE: peptide
119     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
121 Arg Val Gly Gln Cys Thr Asp Ser Asp Val Arg Arg Pro Trp Ala Arg
122 1               5                   10                  15
123 Ser Cys Ala His Gln Gly Cys Gly Ala Gly Thr Arg Asn Ser His Gly
124             20                  25                  30
125 Cys Ile Thr Arg Pro Leu Arg Gln Ala Ser Ala His
126         35                  40
128 (2) INFORMATION FOR SEQ ID NO: 5:
130     (i) SEQUENCE CHARACTERISTICS:
131         (A) LENGTH: 39 amino acids
132         (B) TYPE: amino acid
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RAW SEQUENCE LISTING   DATE: 05/30/2000
PATENT APPLICATION: US/09/079,678A   TIME: 21:32:35

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133        (C) STRANDEDNESS:
134        (D) TOPOLOGY: unknown
136    (ii) MOLECULE TYPE: peptide
138    (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
140  Ser His Ser Gly Gly Met Asn Arg Ala Tyr Gly Asp Val Phe Arg Glu
141  1               5                   10                  15
142  Leu Arg Asp Arg Trp Asn Ala Thr Ser His His Thr Arg Pro Thr Pro
143              20                  25                  30
144  Gln Leu Pro Arg Gly Pro Asn
145              35
147 (2) INFORMATION FOR SEQ ID NO: 6:
149      (i) SEQUENCE CHARACTERISTICS:
150          (A) LENGTH: 41 amino acids
151          (B) TYPE: amino acid
152          (C) STRANDEDNESS:
153          (D) TOPOLOGY: unknown
155     (ii) MOLECULE TYPE: peptide
157     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
159  Ser Pro Cys Gly Gly Ser Trp Gly Arg Phe Met Gln Gly Gly Leu Phe
160  1               5                   10                  15
161  Gly Gly Arg Thr Asp Gly Cys Gly Ala His Arg Asn Arg Thr Ser Ala
162              20                  25                  30
163  Ser Leu Glu Pro Pro Ser Ser Asp Tyr
164              35                  40
166 (2) INFORMATION FOR SEQ ID NO: 7:
168      (i) SEQUENCE CHARACTERISTICS:
169          (A) LENGTH: 39 amino acids
170          (B) TYPE: amino acid
171          (C) STRANDEDNESS:
172          (D) TOPOLOGY: unknown
174     (ii) MOLECULE TYPE: peptide
176     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
178  Arg Gly Ala Ala Asp Gln Arg Arg Gly Trp Ser Glu Asn Leu Gly Leu
179  1               5                   10                  15
180  Pro Arg Val Gly Trp Asp Ala Ile Ala His Asn Ser Tyr Thr Phe Thr
181              20                  25                  30
182  Ser Arg Arg Pro Arg Pro Pro
183              35
185 (2) INFORMATION FOR SEQ ID NO: 8:
187      (i) SEQUENCE CHARACTERISTICS:
188          (A) LENGTH: 44 amino acids
189          (B) TYPE: amino acid
190          (C) STRANDEDNESS:
191          (D) TOPOLOGY: unknown
193     (ii) MOLECULE TYPE: peptide
195     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
197  Ser Gly Gly Glu Val Ser Ser Trp Gly Arg Val Asn Asp Leu Cys Ala
198  1               5                   10                  15
199  Arg Val Ser Trp Thr Gly Cys Gly Thr Ala Arg Ser Ala Arg Thr Asp
```

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        RAW SEQUENCE LISTING            DATE: 05/30/2000
        PATENT APPLICATION:  US/09/079,678A    TIME: 21:32:35

        Input Set : A:\Pto.amc
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200         20              25              30
201  Asn Lys Gly Phe Leu Pro Lys His Ser Ser Leu Arg
202         35              40
204 (2) INFORMATION FOR SEQ ID NO: 9:
206     (i) SEQUENCE CHARACTERISTICS:
207          (A) LENGTH: 44 amino acids
208          (B) TYPE: amino acid
209          (C) STRANDEDNESS:
210          (D) TOPOLOGY: unknown
212     (ii) MOLECULE TYPE: peptide
214     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
216  Ser Asp Ser Asp Gly Asp His Tyr Gly Leu Arg Gly Gly Val Arg Cys
217  1             5              10              15
218  Ser Leu Arg Asp Arg Gly Cys Gly Leu Ala Leu Ser Thr Val His Ala
219         20              25              30
220  Gly Pro Pro Ser Phe Tyr Pro Lys Leu Ser Ser Pro
221         35              40
223 (2) INFORMATION FOR SEQ ID NO: 10:
225     (i) SEQUENCE CHARACTERISTICS:
226          (A) LENGTH: 39 amino acids
227          (B) TYPE: amino acid
228          (C) STRANDEDNESS:
229          (D) TOPOLOGY: unknown
231     (ii) MOLECULE TYPE: peptide
233     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
235  Arg Ser Leu Gly Asn Tyr Gly Val Thr Gly Thr Val Asp Val Thr Val
236  1             5              10              15
237  Leu Pro Met Pro Gly His Ala Asn His Leu Gly Val Ser Ser Ala Ser
238         20              25              30
239  Ser Ser Asp Pro Pro Arg Arg
240         35
242 (2) INFORMATION FOR SEQ ID NO: 11:
244     (i) SEQUENCE CHARACTERISTICS:
245          (A) LENGTH: 38 amino acids
246          (B) TYPE: amino acid
247          (C) STRANDEDNESS:
248          (D) TOPOLOGY: unknown
250     (ii) MOLECULE TYPE: peptide
252     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
254  Arg Thr Thr Thr Ala Lys Gly Cys Leu Leu Gly Ser Phe Gly Val Leu
255  1             5              10              15
256  Ser Gly Cys Ser Phe Thr Pro Thr Ser Pro Pro Pro His Leu Gly Tyr
257         20              25              30
258  Pro Pro His Ser Val Asn
259         35
261 (2) INFORMATION FOR SEQ ID NO: 12:
263     (i) SEQUENCE CHARACTERISTICS:
264          (A) LENGTH: 39 amino acids
265          (B) TYPE: amino acid
```

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                    RAW SEQUENCE LISTING              DATE: 05/30/2000
                    PATENT APPLICATION:  US/09/079,678A      TIME: 21:32:35

                    Input Set : A:\Pto.amc
                    Output Set: N:\CRF3\05262000\I079678A.raw

266              (C) STRANDEDNESS:
267              (D) TOPOLOGY: unknown
269         (ii) MOLECULE TYPE: peptide
271         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
273     Ser Pro Lys Leu Ser Ser Val Gly Val Met Thr Lys Val Thr Glu Leu
274     1               5              10               15
275     Pro Thr Glu Gly Pro Asn Ala Ile Ser Ile Pro Ile Ser Ala Thr Leu
276                 20                  25                  30
277     Gly Pro Arg Asn Pro Leu Arg
278             35
280  (2) INFORMATION FOR SEQ ID NO: 13:
282         (i) SEQUENCE CHARACTERISTICS:
283              (A) LENGTH: 39 amino acids
284              (B) TYPE: amino acid
285              (C) STRANDEDNESS:
286              (D) TOPOLOGY: unknown
288         (ii) MOLECULE TYPE: peptide
290         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
292     Arg Trp Cys Gly Ala Glu Leu Cys Asn Ser Val Thr Lys Lys Phe Arg
293     1               5              10               15
294     Pro Gly Trp Arg Asp His Ala Asn Pro Ser Thr His His Arg Thr Pro
295                 20                  25                  30
296     Pro Pro Ser Gln Ser Ser Pro
297             35
299  (2) INFORMATION FOR SEQ ID NO: 14:
301         (i) SEQUENCE CHARACTERISTICS:
302              (A) LENGTH: 44 amino acids
303              (B) TYPE: amino acid
304              (C) STRANDEDNESS:
305              (D) TOPOLOGY: unknown
307         (ii) MOLECULE TYPE: peptide
309         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
311     Arg Trp Cys Gly Ala Asp Asp Pro Cys Gly Ala Ser Arg Trp Arg Gly
312     1               5              10               15
313     Gly Asn Ser Leu Phe Gly Cys Gly Leu Arg Cys Ser Ala Ala Gln Ser
314                 20                  25                  30
315     Thr Pro Ser Gly Arg Ile His Ser Thr Ser Thr Ser
316             35                  40
318  (2) INFORMATION FOR SEQ ID NO: 15:
320         (i) SEQUENCE CHARACTERISTICS:
321              (A) LENGTH: 39 amino acids
322              (B) TYPE: amino acid
323              (C) STRANDEDNESS:
324              (D) TOPOLOGY: unknown
326         (ii) MOLECULE TYPE: peptide
328         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
330     Ser Lys Ser Gly Glu Gly Gly Asp Ser Ser Arg Gly Glu Thr Gly Trp
331     1               5              10               15
332     Ala Arg Val Arg Ser His Ala Met Thr Ala Gly Arg Phe Arg Trp Tyr
```

Davenport N

Serial Number: 09/079,678A

ENTERED

PH 11

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically: _____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: _____

☐ Deleted extra, invalid, headings used by an applicant, specifically: _____

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____ .

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically: _____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: _____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☑ Other: Seq 179 - fixed a.a. no

_____

_____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95